LEAP-Agri-project OPTIBOV

Local traditional breeds,

well-adapted to their ecosystems, **will be lost** if crossbreeding or breed replacement are not monitored

OUR AIM

To contribute key information towards improving the performance of traditional cattle breeds while maintaining their biodiversity

HOW?

We want to find DNA markers in the genomes of your animals related to ecosystem adaptation and disease resistance which can be used together with production markers for future genetic improvement

WHAT WE ASK

Your collaboration to **collect biological samples** and record phenotypes such as diseases, production, abnormalities, etc., in a few selected animals over the course of 2 years

WHAT'S IN IT FOR YOU

- You and your animals will be a part of an international scientific research project
- The information records of both your animals and other animals in the study are available for you to consult and compare
- Access to our database through the OPTIBOV app on your mobile, where you can upload your records and check information on your animals
- Useful genomic information for improved animals by selection
- · A reward certificate after 2 years in this research project





EUROPE

North, Finland







Eastern, Northern, Western Finncattle

Center, Netherlands







Dutch Belter Lakenvelder

Groninger Whiteheaded (Blaarkop)

Meuse-Rhine-Yssel

South, Portugal







Barrosã

Mertolenga

Mirandesa

AFRICA

North, Egypt







Baladi: Damietta, Maryuti, Menufi

Center, Uganda

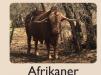




Ankole

East African Zebu

South, South Africa







Nguni

Tuli

TRAITS/INFORMATION OF INTEREST

- Ecosystem/climate conditions
- Temperature/humidity index
- · Availability of water/ food
- Food quality
- Type of food (grazing, supplements, other)
- Disease recording
- Disease resistance
- Parasite infections (e.g. ticks, nematodes)
- Housing (outdoor/indoor)
- Production records
- Calving interval/total number of calves
- · Age at first calf
- Longevity
- Treatments
- Biochemical measurements on blood
- Immunity parameters
- MHC haplotypes

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